

## CLAIM AMENDMENTS

1. (Currently Amended) ~~Fermentation~~ A fermentation process for the preparation of ~~L-amino acids~~ a desired L-amino acid selected from the group consisting of L-threonine, L-isoleucine, L-valine, and L-lysine, especially L-threonine, wherein the following steps are carried out:

a) ~~fermentation of the microorganisms of the family Enterobacteriaceae of an *E.coli* strain in a fermentation broth for producing the desired L-amino acid, in which microorganisms at least the *pekA* gene or nucleotide sequences coding thereof are attenuated and, in particular, switched off wherein the endogenous gene encoding phosphoenolpyruvate (PEP) carboxykinase (*pckA* gene) of *E.coli* is attenuated,~~

b) ~~enrichment of the L-amino acid in the medium or in the bacterial cells~~ concentration of the fermentation broth to eliminate water and increase the concentration of said L-amino acids in the broth and *E.coli* , and

c) ~~isolation of the L-amino acid, constituents of the fermentation broth and the biomass in its entirety or portions thereof optionally being isolated as a solid product together with the L-amino acid.~~

2. (Currently Amended) ~~Process~~ The process according to claim 1, wherein ~~microorganisms are used in which~~ other genes of the biosynthetic pathway of the desired L-amino acid of *E. coli* are additionally amplified.

3. Canceled

4. (Currently Amended) ~~Process~~ The process according to claim 1, wherein the expression of the ~~polynucleotide(s) coding for the *pckA* gene~~ is attenuated and, in particular, ~~switched off.~~

5. (Currently Amended) ~~Process~~ The process according to claim 1, wherein the regulatory and/or catalytic properties of the ~~polypeptide (enzyme protein) coded for by the polynucleotide *pekA*~~ polypeptide encoded by the *pckA* gene are reduced.

6. (Currently Amended) ~~Process~~ The process according to claim 1, wherein ~~microorganisms of the family Enterobacteriaceae in which~~ one or more *E.coli* genes selected from the group ~~comprising~~ consisting of:

6.1 the thrABC operon coding for aspartate kinase, homoserine dehydrogenase, homoserine kinase and threonine synthase,

6.2 (a) the pyc gene coding for pyruvate carboxylase,

6.3 (b) the pps gene coding for phosphoenolpyruvate synthase,

6.4 (c) the ppc gene coding for phosphoenolpyruvate carboxylase,

6.5 (d) the pntA and pntB genes coding for transhydrogenase,

6.6 (e) the rhtB gene for homoserine resistance, ~~and~~

6.7 (f) the rhtC gene for threonine resistance, and

6.8 (g) the gdhA gene coding for glutamate dehydrogenase

~~are simultaneously amplified and, in particular, overexpressed are fermented~~ during fermentation for the preparation of said L-amino acids.

7. (Currently Amended) ~~Process~~ The process according to claim 1, wherein ~~microorganisms of the family Enterobacteriaceae in which~~ one or more *E.coli* genes selected from the group ~~comprising~~ consisting of:

7.1 (a) the tdh gene coding for threonine dehydrogenase,

7.2 (b) the mdh gene coding for malate dehydrogenase,

7.3 (c) the gene product of the open reading frame (orf) yjfA, and

7.4 (d) the gene product of the open reading frame (orf) ytfP,

~~are attenuated and, in particular, switched off, or the expression is reduced, are fermented~~ during fermentation for the preparation of said L-amino acids.

8. (Canceled)

9. (Currently Amended) ~~Process~~ The process according to claims 1 to ~~8~~ 7, wherein L-isoleucine, L-valine, L-lysine, or L-threonine is prepared.

10-27. (Canceled)